

1617
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Intellectual Property Law Group PLLC



September 22, 2003

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To whom it may concern:

We herewith return the enclosed Amendment Checklist papers and attached claims, which appear to have been inadvertently placed within the Office Action sent to our office on September 15, 2003 for U.S. Patent Application No. 08/486,867.

These Checklists, referring to U.S. Application Nos. 10/077,588 and 09/697,123, do not correspond to the application to which the Office Action referred, nor are they applications being prosecuted by our firm.

If you have any questions or require additional information, please call.

Sincerely,
SEED Intellectual Property Law Group PLLC

Linda Povinelli
For Jeffrey C. Pepe

Enclosures: As noted

(JCP/lmp)

418151



1. (Amended) A DNA fragment which has a sequence selected from the group consisting of SEQ ID NOS:1 to 4 and 6 to 24.

2. (Amended) A method for identifying the species or subspecies of a mycobacterial strain comprising the steps of:

- a) digesting a DNA fragment which has a sequence selected from the group consisting of SEQ ID NO:1 to SEQ ID NO:24 with at least one restriction enzyme selected from the group consisting of *HaeIII*, *MspI*, *Sau3AI*, and *BstEII* to obtain a first DNA fragment length polymorphism pattern;
- b) isolating a DNA fragment from the mycobacterial strain to be identified;
- c) amplifying *rpoB* region of the DNA fragment isolated in step (b), said amplification being performed by using a primer of SEQ ID NO: 15 or SEQ ID NO: 26;
- d) digesting the DNA fragment amplified in step c) with the at least one restriction enzyme employed in step a) to obtain a second DNA fragment length polymorphism pattern;
- e) comparing the first DNA fragment length polymorphism pattern obtained in step a) with the second DNA fragment length polymorphism pattern obtained in step d), thereby identifying the species or subspecies of a mycobacterial strain.

3. (Amended) A method of claim 2, wherein said first and second DNA fragment length polymorphism by electrophoresis.

5. (Amended) A method of claim 2, wherein said mycobacterial strain is selected from the group consisting of *M. tuberculosis*, *M. avium*, *M. abscessus*, *M. flavescens*, *M. africanum*, *M. bovis*, *M. chelonae*, *M. celatum*, *M. fortuitum*, *M. gordonae*, *M. gastr*, *M. haemophilum*, *M. intracellulare*, *M. kansasii*, *M. mageritensis*, *M. marinum*, *M. szulgai*, *M. terrae*, *M. scrofulaceum*, *M. ulcerans*, and *M. xenopi*.

IN THE DRAWINGS

Corrected Figs. 6a-6b are attached.